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OM protein - protein search, using sw model
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Listing first 45 summaries
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/cgn2_6/ptodata/2/iaa/FCTUS_COMB.pep:*
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39.701 Million cell updates/sec
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ALIGNMENTS

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; Sequence 146_Application US/08940095
; Patent_MO. 6004925
; GENERAS_INSCRMATION:
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; TOPOLOGY: 1; MOLECULE TYPE: US-08-940-095-146
                                                                                                                                             NAME: COTUZZÍ, LAUTA A
REGISTRATION NUMBER: 30,742
REFERRNCE/DOCKET NUMBER: 0991
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEPAX: 650-493-4935
TELEX: 66141 PENNIE
                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dufourcq, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
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APPLICATION NUMBER:
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                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Penute a Long STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 29-SE
                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
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Cornut, Isabelle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         IBM Compatible
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V: 514
                      No. 6004925e
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Query Match

DB 3; Length 22;

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 00-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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                                                                                                                                 LOCATION: 1...22
OTHER INFORMATION: All amino acids are in the D-configuration
                                                                                                                                                                                                                                 LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                           TELEPHONE: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM CO
OPERATING SYSTEM:
SOFTWARE: FastSE
                                                                                                                                                                        NAME/KEY: Other
             1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                      TOPOLOGY:
1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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o. 6004925
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                                                         Conservative
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Cornut, Isabelle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pennie & Edmonds LLP
                                                                                                                                                                                                     No. 6004925e
                                                                   100.0%; Score 105; DB 3 100.0%; Pred. No. 2e-08;
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100.0%; Pred. No. ...
tive 0; Mismatch
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                                                                                                                                                                                                                                                                                           174:
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                                                       Mismatches
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                                                                              DB 3; Length 22;
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                                                    Indels
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US-08-940-093-174
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                                                                                                                                                                             RESULT
                                                                                                                           Patent No.
                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-940-093-146
                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                    GENERAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 146, Application US/08940093 Patent No. 6037323
                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
            TITLE OF INVENTION:
 NUMBER OF SEQUENCES:
                                           APPLICANT:
                                                       APPLICANT:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 00 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: APOLIPOPROTEIN A-I ACONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 650-493-5556
                                                                                                                                                                                                                  1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                 1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Coruzzi, Laura A REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 29-SEI
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SOFTWARE: FastSE
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                                                                                                                                            124 Application US/08940093
                                                                                                                                                                                                                                                                                22;
                                                                                                                                 6037323
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                                                                                                                                                                                                                                                                                                                                                                                                                             22 amino acids
                                        Metz, Gunther
                                                    Buttner, Klaus
Cornut, Isabelle
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1155 Avenue of the Americas
                                                                                     Sekul, Renate
                                                                                                 Dasseux,
                                                                                                                                                                                                                                                                              Conservative
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Cornut, Isabelle
Metz, Gunther
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                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    650-493-4935
                                                                                                                                                                                                                                                                                                                                                                   No. 6037323e
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         AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                        APOLIPOPROTEIN A-I AGONISTS
                                                                                                   Jean-Louis
                                                                                                                                                                                                                                                                                      100.0%; Score 105; DB 3 100.0%; Pred. No. 2e-08;
258
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

E: Pennie & Edmonds LLP 1155 Avenue of the Americas

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LOCATION: 1...22
OTHER INFORMATION: All amino acids are in the D-configuration
US-08-940-093-174
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CURRENT APPLICATION NUMBER: US/08/940,
APPLICATION NUMBER: US/08/940,
                                                                                                                                                                                                                                                              В
                                                                                                                                                                                             ; sequence 146,
; patent No. 604
                                                                                                                                                                                                                   US-08-940-096-146
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 435
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
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CITY: New York
                                                                                                                                                                                                                              RESULT
                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: No. 6037323e
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                      GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                   APPLICANT:
                                                                                                                                                     APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
                                                                                                            TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                        Local Similarity
les 22; Conserv
                                                                                          NUMBER OF SEQUENCES: 2:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                               1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                   1 PVLELFENLLERLLDALQKKLK 22
          ZIP: 10030 FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                       STREET:
                                                                                  ADDRESSEE:
                                                  STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Coruzzi
                                         COUNTRY :
COMPUTER:
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                                                                                                                                                                                        INFORMATION:
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                                10036-2811
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                                                               New York
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N: 435
                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                              Application US/08940096
                                                       ΝY
                                                                          1155 Avenue of the Americas
                                                                                                                                                  Buttner, Klaus
Cornut, Isabelle
                                             USA
                                                                                    pennie & Edmonds LLP
  IBM Compatible
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                                                                                                                                                                                                                                                                                                                              100.0%;
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pred. No. 2e-08;
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                                                                                                                                                                                                                                                                                                                                               DB 3; Length 22;
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MOLECULE TYPE: NO. 6046166e
US-08-940-096-146
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SOFTWARE: FASTSEO VETSION 2.0
CURRENT APPLICATION DATA:
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Patent No. 6046166
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TELEX: 66141 PENNIE
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 146:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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RE
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Bekul, Renate
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
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Local Similarity 100.0%; Pred. No. 2
Local Similarity 0; Mismatchen
hes 22; Conservative 0; Mismatchen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLELFENLLERLLDALQKKLK 22
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1155 AV
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                    CLASSIFICATION: 530
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
NAME: COTUZZI, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 005
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM CON OPERATING SYSTEM:
                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 29-SE
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29-SEP-1997
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                                                                                                                                                                                                                                        umber: US/08/940,096
29-Sep-1997
N: E77
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                                                              009196-0005-999
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INFORMATION FOR SEQUENCE CHARACTERISTICS:
LENGTH: 22 millo acids
STRANDEDNESS: Single
MOLECHY 5 Lines
                                                                                REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 0091
TELEPHONE: 500-493-935
TELEPHONE: 650-493-935
      MOLECULE TYPE:
                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-465-719-146
                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                     SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
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LOCATION: 1...2

OTHER INFORMATION: All amino acids are in the D-configuration
                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent,
                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                               OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
                                                                                                                                                                                                                                                               COMPUTER:
                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 10036-2811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
INFORMATION FOR SEO ID NO: 174:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: No. 6046166e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PVLELFENLLERLLDALQKKLK 22
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Local Similarity 100.0%; Score 105; DB 3
les 22; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        -146-Application US/09465719
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                   New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION
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Cornut, Isabelle
Metz, Gunther
No. 6265377e
                                                                                                                                                                                                                                                      IBM Compatible
                                                                                                                                                                                                                                                                   Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                        009196-0006-999
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    Matches
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US-09-465-719-174
                                                                               Query Match
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TELEFAX: 650-493-5556

TELEX: 66141 PENNIE
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
STRANDEDNESS: single
TOPOLOGY. Single
                                                                                                                                                          MOLECULE TYPE: No. 6265377e
1 PVLELFENLLERLLDALQKKLK 22
                    1 PVLELFENLLERLLDALQKKLK 22
                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-465-719-174
                                                                                        OTHER INFORMATION: All amino acids are in the D-configuration
                                                                                                                                                                                                                                                                      NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 000
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 174, ; Patent No. 62
                                                                                                                                NAME/KEY: Other LOCATION: 1...2;
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/94
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskett
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ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
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CITY: New York
CTATE: NY
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APPLICANT:
APPLICANT:
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APPLICANT:
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Best Local (
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TITLE OF INVENTION: AND THEIR USE TO TREAT DYST.
                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
SOFTWARE: Fast SEC
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les 22; Conserv
                                              Conservative
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                                                                                                                                                                           linear
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Cornut, Isabelle
Metz, Gunther
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                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM Compatible
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Sekul, Renate
                                                                                                                                                                                   single
                                             100.0%; Score 105; DB 4
100.0%; Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                        08/940,093
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100.0%; Pred. No. 2e-08;
tive 0; Mismatches
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                                      Mismatches
                                                        DB 4;
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                                                  Length 22;
                              Indels
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                        0; Gaps
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US-08-940-095-150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dasseux, Jean
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabell
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
                                                                                                                                                                                                                                                                             TOPOLOGY: 11;
; MOLECULE TYPE:
; OS-940-095-150
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CORRESPONDENCE ADDRESS:
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APPLICANT: DUCOURCY, Jean
APPLICANT: DUCOURCY APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
TITLE OF INVENTION: 258
                                                                                  ; Sequence 159, Application US/08940095; Patent No. 6004925
                                                                                                                            RESULT
                                                                                                                                                                                                                                                                                                                                                                         TELEPAX: 650-493-5556
TELEX: 66141 PENNIE
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 150:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 29-SEP-1
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
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CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: COTUZZI, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                        APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
                                                                                                                                                                                                                   Local Similarity
es 21; Conserv
                                                                                                                                                                   1 PVLELFENLLERLLDALOKKLK 22
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Cornut, Isabelle
                                                                                                                                                                                                                                                                                                                                          amino acid
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                                                                                                                                                                                                                                                                                                                                                    22 amino acids
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                                                                                                                                                                                                                         conservative
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                 Sekul, Renate
Buttner, Klaus
Cornut, Isabelle
Metz, Gunther
                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-SEP-1997
       Dufourcq, Jean
                                                                       passeux,
                                                                                                                                                                                                                                                                                                     No. 6004925e
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                                                                                                                                                                                                                                                                                                                             single
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                                                                                                                                                                                                                                             97.1%;
                                                                         Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                         009196-0004-999
                                                                                                                                                                                                                                            Score 102; DB 3;
Pred. NO. 5.2e-08;
                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                               DB 3; Length 22;
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MOLECULE TYPE: NO. 6004925e
US-08-940-095-159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 650-493-5556
TELEX: 66141 PENNIE
TELEX: 66141 PINO:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FASTSEO Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
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CITY: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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NAME: COTUZZI, LAULG 30,742
REGISTRATION NUMBER: 00:
REFERENCE/DOCKET N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         APPLICANT: Dasseux, Jean-L.
APPLICANT: Sekul, Renate
APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabell.
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
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                                                                                                                                                                                                                                                                                      APPLICANT: Dufourcy, Jean TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 21; Conserv
                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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                                                                                                                              CITY: New York
STATE: NY
                                                                                                                                                                                     ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                   COUNTRY:
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                                                                                 10036-2811
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Cornut, Isabelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-SEP-1997
                                                                                                                     USA
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AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97.1%; Score 102; DB 3; Length 22; 95.5%; pred No. 5.2e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jean-Louis
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CURRENT APPLICATION DATA:

FastSEQ Version 2.0

US/08/940,095

SOFTWARE: OPERATING SYSTEM:

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US-08-940-095-179
                                             ATTORNEY/AGENT INFORMATION:
                                                                      CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 179,
Patent No. 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ŪS-08-940-095-169
                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,095
FILING DATE: 29-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 650-493-5556
INFORMATION FOR SEQ ID NO: 16
SEQUENCE CHARACTERISTICS: 16
LENGTH: 2 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                APPLICANT: Sekul, Neille APPLICANT: Buttner, Klaus APPLICANT: Buttner, Klaus APPLICANT: Cornut, Isabelle APPLICANT: Metz, Gunther APPLICANT: Dufourcq, Jean TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
            REFERENCE/DOCKET NUMBER: 00
                      NAME: COruzzi, Laura A REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                           COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 2.0
                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PVLELFENLLEKLLDALQKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PVLELFENLLERLLDALQKKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 650-450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: COTUZZI, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                 10036-2811
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/0
FILING DATE: 29-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 /9, Application US/08940095
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                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No. 6004925e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97.1%; Score 102; DB 3; 95.5%; Pred: No. 5.2e-08;
009196-0004-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   009196-0004-999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 22;
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US-08-940-095-188
                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEX: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 188:
LENGTH: 22 amino acids
TYPE: amino acids
TYPE: amino acids
          STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,0:
EILING DATE: 29-SEP-1997
PRIOR APPLICATION: 514
ADDITIONATION: DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 650-493-5556
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS: 2 amino acids
TYPE: amino acids
STRANDEDNESS: single
MOLECULE TYPE: NO. 6004925,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-940-095-188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 188, Applicate Patent No. 6004925
GENERAL INFORMATION:
                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Sekul, Renate
APPLICANT: Sekul, Renate
APPLICANT: Ornut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
NUMBER OF SEQUENCES: 258
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                    ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                        NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
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                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM CO
OPERATING SYSTEM:
SOFTWARE: FastSE
                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 10036-2811
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TELEPHONE: 650-493-4935
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                            linear
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      No. 6004925e
                                                                                                                                                                                                                                                                                                                                        IBM Compatible
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                                single
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    Mismatches

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; Sequence :
; Patent NO.
; GENERAL IN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 150, Application US/08940093 patent No. 6037323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                  STRANDEDNESS: single
TOPOLOGY: linear
TOPOLOGY: NO. 6037323e
MOLECULE TYPE: NO. 6037323e
US-08-940-093-150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dasseux, Jean-
APPLICANT: Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
                                                                         В
                       US-08-940-093-159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                      RESULT
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 150:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                        Matches
                                                                                                                                    Query Match
Best Local Similarity
Sequence 159, Application US/08940093 patent No. 6037323
                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                   NAME: GOTUZZI, LAURA 30,742
REGISTRATION NUMBER: 30,742
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
650-493-5556
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                                                                                                                                    LENGTH:
                                                                         10036-2811
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1155 Avenue of the Americas
                                                                                                                               21;
                                                                                                                                                                                                                                   amino acids
                                                                                                                                                                                                                                                                                        66141 PENNIE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM Compatible
                                                                                                                               conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1997
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95.5%;
                                                                                                                                                                                                                                                                                                                                                  , Laura A 30,742 00'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jean-Louis
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                                                                                                                                              97.1%;
95.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                 009196-0006-999
                                                                                                                                                 Score 102; DB 3; Length 22; pred. No. 5.2e-08;
                                                                                                                                        Mismatches
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                                                                                                                                                      0;
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GENERAL INFORMATION:
                                                                                                                                                                   US-08-940-093-159
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APPLICANT: Buttner, Klaus
APPLICANT: Cornut, Isabelle
APPLICANT: Metz, Gunther
APPLICANT: Metz, Gunther
                                                              Дb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
Search completed: November 20, 2001, 09:43:25 Job time: 189 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                        TELEX: 66141 PENNIE
INFORMATION FOR SEO ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/940,
                                                                                                          Matches
                                                                                                                       Best Local Similarity
                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
CITY:
                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 00
REFERENCE/DOCKET NUMBER: 100
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP:
                                                                                                                                                                                    STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: No. 6037323e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                  FILING DATE
                                                                                                                                                                                                                                                                                                                              NAME: COTUZZI, Laure
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
                                                                                                                                                                                                                                    LENGTH:
                                                               1 PVLELFENLLERLLDALOKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10036-2811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
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Sekul, Renate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ΝY
                                                                                                                 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Pennie & Edmonds LLP
1155 Avenue of the Americas
                                                                                                                                                                                                                       H: 22 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Diskette
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                US/08/940,093
29-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jean-Louis
                                                                                                                                                                                                                                                                                                                                                   Laura A
                                                                                                                                   97.1%;
95.5%;
                                                                                                                                                                                                                                                                                                                                           30,742
                                                                                                                                                                                                                                                                     159:
                                                                                                                                                                                                                                                                                                                               009196-0006-999
                                                                                                                                    Score 102; DB 3;
pred No. 5.2e-08;
                                                                                                                             Mismatches
                                                                                                                                                          Length 22;
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Page 8

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OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run on:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minimum DB seq length: 0 Maximum DB seq length: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0% Maximum Match 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Database :
A_Geneseq_0601:*
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2: /SIDSB/gcgdata,
3: /SIDSB/gcgdata,
4: /SIDSB/gcgdata,
4: /SIDSB/gcgdata,
5: /SIDSB/gcgdata
6: /SIDSB/gcgdata
6: /SIDSB/gcgdata
7: /SIDSB/gcgdata
9: /SIDSB/gcgdata
10: /SIDSB/gcgdata
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105
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Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                        /SIDS8/gcgdata/geneseq/geneseqp/AA1980.DAT:*
/SIDS8/gcgdata/geneseq/geneseqp/AA1981.DAT:*
/SIDS8/gcgdata/geneseq/geneseqp/AA1982.DAT:*
/SIDS8/gcgdata/geneseq/geneseqp/AA1983.DAT:*
/SIDS8/gcgdata/geneseq/geneseqp/AA1984.DAT:*
/SIDS8/gcgdata/geneseq/geneseqp/AA1984.DAT:*
                                                                                                                                                                                                                                                                                                                                         /SIDSB/gcgdata/geneseq/geneseqp/AA1986.DAT:*
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

1 2 3 4 4 6 5 7 7 10 11	Result
105 105 105 105 105 105 105 105 102 102 102	Score
100.0 100.0 100.0 100.0 100.0 100.0 100.0 100.0 97.1 97.1	% Query Match L
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20 AAY19107 20 AAY18853 20 AAY18825 20 AAY18825 20 AAY18590 20 AAY19361 20 AAY19333 20 AAY19333 20 AAY19333 20 AAY19333	ID
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96.2 95.2	96.2	96.2	96.2	96.2	96.5	96.2	96.2	96.2	96.2	96.2	96.2	96.2	96.2	96.2	96.2	97.1	9/.1	97.1	97.1	97.1	97.1	97.1	97. L	97.1	97.1	97.1	97.1	9/.1	97.1	9/.	97.1	97.1	
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AAY 1935	AAY193	Ą		ΑA	ΑA	ΑA	ΑA	AA	AA	AA	A												AAV18595	SERIAN	201983	AA1 1000	AA11001	2001001	AAY1909	8091908	AAI 1 31 6 1 AAV 1 9 1 0 2	AAYLYLL	
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ALIGNMENTS

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AAY19107
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                                                                                                                                                                                                                                                                                                                                                         AAY19107 standard; Peptide; 22 AA.
                                                                                                                                                                                                                              Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                      Lecithin:cholesterol acyltransferase activation exhibiting peptide #174.
                                                                                                                                                                                                                                                                                                               09-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                      AAY19107;
                                                                                                                                                                                                                      septic shock.
                                                                                                                                                                                                Synthetic.
                                 (DASS/)
(METZ/)
                                                                                                                                        08-APR-1999.
                                                                                                                                                              WO9916458-A1
                                                                                                                                                                                    Homo sapiens
                                                          (BUTT/) BUTTNER K. (CORN/) CORNUT I.
                                                                                           29-SEP-1997;
                                                                                                                 28-SEP-1998;
Buttner K,
                       (SEKU/) SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                   DASSEUX J.
METZ G.
  Cornut I, Dasseux J,
                                                                                            97US-0940096.
                                                                                                                    98WO-US20326.
       Metz G,
         Sekul R;
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AAY19079
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Claim 18; Page 115; 254pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                            Peptide agonists of apolipoprotein A-I
                                                          WPI; 1999-277034/23
                                                                                 Buttner K, Cornut I,
                                                                                                                                                (BUTT/) BUTTNER K. (CORN/) CORNUT I.
                                                                                                                                                                                                                                      08-APR-1999
                                                                                                                                        (DASS/)
                                                                                                                                                                                     29-SEP-1997;
                                                                                                                                                                                                             28-SEP-1998;
                                                                                                                                                                                                                                                              WO9916458-A1
                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                         human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid dyslipidemia, specifically hypercholesterolaemia, associated with disease, are used to treat or prevent diseases associated with disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or treating septic shock. When labeled, (A) can also be used diagnostically is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. ANY18937 represent lecithin:cholesterol transferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                              Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia;
                                                                                                                                                                                                                                                                                                                                                                                    Lecithin:cholesterol acyltransferase activation exhibiting peptide #146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY19079 standard; Peptide; 22
                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY19079;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 116; 254pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-277034/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide agonists of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PVLELFENLLERLLDALOKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                          SEKUL R.
                                                                                                                                 DASSEUX J.
                                                                                                                      METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                 97US-0940096
                                                                                                                                                                                                          98WO-US20326.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 105; DB 20;
100.0%; Pred. No. 2.3e-08;
tive 0; Mismatches 0;
                                                                                Dasseux J,
                                                                              Metz
                                                                         e,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
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 dyslipidemia, specifically hypercholesterolaemia,
          The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 14-22 residue peptide, or analog, that forms an amphipathic alpha-hellx in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                     Example; Page 113; 152pp; English.
                                                                                                                  Peptide agonists of apolipoprotein A-I
                                                                                                                                                   WPI; 1999-277031/23.
                                                                                                                                                                                 Buttner K,
                                                                                                                                                                                                                                    (CORN/) CORNUT I. (DASS/) DASSEUX J.
                                                                                                                                                                                                              (SEKU/) SEKUL R.
                                                                                                                                                                                                                              (METZ/)
                                                                                                                                                                                                                                                                 (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                                                               29-SEP-1997;
                                                                                                                                                                                                                                                                                                                          28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                       08-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                W09916408-A2.
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lecithin:cholesterol acyltransferase activation exhibiting peptide #174.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY18853 standard; Peptide; 22 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids (A), and their lipid complexes, are used to treat or prevent diseases associated with disease, are therosclerosis, restenosis, HDL (high density lipoprotein) or treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at AAY19187 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 pvlelfenllerlldalqkkik 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ω
                                                                                                                                                                                                                        METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                              Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                            97US-0940093
                                                                                                                                                                                                                                                                                                                      98WO-US20328.
                                                                                                                                                                           Dasseux J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 105; DB 20;
Pred. No. 2.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                           Metz G,
                                                                                                                                                                        Sekul R;
cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT
AAY18825
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY18680 to AAY18933 represent lecithin cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY18825 standard; Peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-APR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY18825;
                                 apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that
                                                                                                                         The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 14-22 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               septic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lecithin: cholesterol acyltransferase activation exhibiting peptide #146.
                                                                                       dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or
                                                                                                                                                                                                                                                                                        WPI; 1999-277031/23.
                                                                                                                                                                                                                                                                                                                          Buttner K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-SEP-1998;
e.g. atherosclerotic streaks, and to raise antibodies.
                                                                                                                                                                                                                 Example; Page 112; 152pp; English.
                                                                                                                                                                                                                                                    Peptide agonists of apolipoprotein A-I
                    is involved in retrograde cholesterol transport, also to image HDL at
                                                                                                                                                                                                                                                                                                                                                                                             (DASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                (CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                               (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 pvlelfenllerildalqkkik 22
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METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                                CORNUT I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                            Dasseux J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 105; DB 20;
Pred. No. 2.3e-08;
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                                                                                                                                                                                                                                                                                                                            Metz G,
                                                                                                                                                                                                                                                                                                                             Sekul R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
         AAY18680 to
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RESULT
AAY18590
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY18933 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                      are useful for gene therapy, or prevention, of diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) and apoA-I deficiency, hypertriglyceridemia and metabolic syndrome, also to treat endotoxemia (septic shock). Host cells containing (A) can also be used to study the role of apoA-I in lipid metabolism. (B) can be used diagnostically, e.g. to measure serum HDL (particularly its subpopulation involved in retrograde cholesterol transport) and for simaging the circulatory system or HDL accumulations at fatty streaks.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene therapy; apolipoprotein A-I; agonist; dyslipidemic disorder; ApoA-I; cardiovascular disease; atherosclerosis; restenosis; LCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lecithin:cholesterol acyltransferase activation exhibiting peptide #174.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY18590 standard; Peptide;
                                                                                                                                                                         The present invention describes a nucleic acid (A) encoding an apolipoprotein A-I (apoA-I) agonist (B) that is a peptide, or and which forms an amphipathic alpha-helix in presence of lipids. (A) optionally as a complex with lipids, and host cells that contain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hyperlipidemia; septic shock; lecithin:cholesterol acyltransferase
                                                                                                                                                                                                                                                                                                                                        Sekul R;
                                                                                                                                                                                                                                                                                                                                                         Buttner K, Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                              (CORN/) CORNUT I.
(DASS/) DASSEUX J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9916409-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                              Nucleic acid encoding apolipoprotein A-I agonist peptides
                                                                                                                                                                                                                                                                                                              WPI; 1999-254921/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                    Example; Page 178; 232pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     (SEKU/) SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                                                 (DUFO/)
                             The present sequence represents a peptide from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                     (METZ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                                                                   METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                                   DUFOURCQ J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-0940136
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                                                                                                                                                                                                                                                                                                                                                           Dasseux J, Dufourcq J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 105; DB 20;
Pred. No. 2.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                              Metz
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                                                                                                                                                                                                             or analog,
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                                                                                                                                                                                               (A),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                Matches
                                      Query Match
Best Local
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                                                                                                                                                                                                  The present invention describes a nucleic acid (A) encoding an apolipoprotein A-I (apoA-I) agonist (B) that is a peptide, or analog, which forms an amphipathic alpha-helix in presence of lipids. (A), optionally as a complex with lipids, and host cells that contain (A), are useful for gene therapy, or prevention, of diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) and apoA-I deficiency, hypertriglyceridemia and metabolic syndrome, also to treat to study the role of apoA-I in lipid metabolism. (B) can also be used diagnostically, e.g. to measure serum HDL (particularly its subpopulation involved in retrograde cholesterol transport) and for imaging the circulatory system or HDL accumulations at fatty streaks.
                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 35; Page 170; 232pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid encoding apolipoprotein A-I agonist peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-254921/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Buttner K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9916409-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene therapy; apolipoprotein A-I; agonist; dyslipidemic disorder; ApoA-I; cardiovascular disease; atherosclerosis; restenosis; LCAT; hyperlipidemia; septic shock; lecithin:cholesterol acyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lecithin: cholesterol acyltransferase activation exhibiting peptide #146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-APR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY18562 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DUFO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                 present sequence represents a peptide from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  \vdash
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Φ
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                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORNUT I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DUFOURCQ J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DASSEUX J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          METZ G
                                                                                                                                              22 AA;
          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97US-0940136.
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                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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        0,
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                Score 105; Db 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 105; DB 20;
Pred. No. 2.3e-08;
; Mismatches 0;
        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dufourcq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Metz G;
     Indels
                                                      Length
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                                                         22;
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0;
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Gaps
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0;
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1 PVLELFENLLERLLDALQKKLK 22

Query Match Best Local Matches 2

Local Similarity

Conservative

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100.0%;

Score 105; DB 20; Pred. No. 2.3e-08; Mismatches 0;

Length Indels

22; 0;

Gaps

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RESULT
AAY19361
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                              The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atheroscierosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY19188 to AAY19441 represent lecithin:cholesterol acyltransferase (LCAT) activity
Sequence
                           exhibiting
                                                                                                                                                                                                                              Example; Page 125; 280pp; English
                                                                                                                                                                                                                                                           Peptide agonists of apolipoprotein
                                                                                                                                                                                                                                                                                             WPI; 1999-277035/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                          Sekul R;
                                                                                                                                                                                                                                                                                                                                       Buttner K, Cornut I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-APR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                               (DASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                             (CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                         (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   septic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lecithin:cholesterol acyltransferase activation exhibiting peptide #174.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY19361 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                   METZ/
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                                                                                                                                                                                                                                                                                                                                                                                              DUFOURCQ J
                                                                                                                                                                                                                                                                                                                                                                                                                          CORNUT I.
                                                                                                                                                                                                                                                                                                                                                                                METZ G.
                                                                                                                                                                                                                                                                                                                                                                                                             DASSEUX J.
22
                          core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
ΑA;
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RESULT
AAY19375
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                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY19333;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lecithin:cholesterol acyltransferase activation exhibiting peptide #146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               septic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                             The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with complexes, are used to treat or prevent diseases, cardiovascular dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or aloa-I deficiency; hypertriglyceridemia and metabolic syndrome, also for apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for apoA-I deficiency; hypertriglyceridemia to an also be used diagnostically treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL suboppulation that is involved in retrograde cholesterol transport, also to image HDL at is involved in retrograde cholesterol transport, also to image HDL at is involved in retrograde cholesterol acyltransferase (LCAT) activity anyly441 represent lecitinicholesterol acyltransferase (LCAT) activity anyly441 represent lecitinicholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9916459-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 124; 280pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide agonists of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-277035/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BUTT/) BUTTNER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DUFO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DASS/) DASSEUX J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SEKU/) SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORN/)
                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                        exhibiting
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AAY19375 standard;
                                                                                                          1 PVLELFENLLERLLDALOKKLK 22
                                                                                                                                        Local Similarity
es 22; Conserv
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                                  9
                                                                             DUFOURCQ J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METZ G
                                                                                                                                                                                                                                                    core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                           22 AA;
                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97US-0940095
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   Peptide;
                                                                                                                                                             100.0%;
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                                                                                                                                                 0;
        22 AA.
                                                                                                                                                             Score 105; DB 20;
pred. No. 2.3e-08;
                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Metz G;
                                                                                                                                                      Indels
                                                                                                                                                                                  Length 22;
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В
                                                                                  RESULT
                                                                                                                                                                                       Matches
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                                                                                                                                                                                                         Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin; cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            septic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lecithin:cholesterol acyltransferase activation exhibiting peptide #188.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY19375;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9916459-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-SEP-1998;
                                                                                                                                                                                                                                                                                                     disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically treating septic serum levels of HDL, in particular the HDL subpopulation that to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at is involved in retrograde cholesterol transport, also to image HDL at is involved in retrograde cholesterol acultransferase (LCAT) activity AAV19441 represent lecithin:cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                           complexes, are used to treat or prevent diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid amphipathic alpha-helix in presence of lipids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 126; 280pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide agonists of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-277035/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sekul R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Buttner K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BUTT/)
                                                                                                                                                                                                                                                                                                                                                                                                              dyslipidemia, specifically hypercholesterolaemia, cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SEKU/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DASS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DUFO/)
                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                               exhibiting
14-JUL-1999
                            AAY19346;
                                                         AAY19346 standard; Peptide;
                                                                                                                                                                                                       Local Similarity
                                                                                      10
                                                                                                                              1 PVLELFENLLERLLDALQKKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DASSEUX J
                                                                                                                                                                                            21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEKUL R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BUTTNER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METZ G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DUFOURCQ J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cornut I,
                                                                                                                                                                                                                                                                                             core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                    22 AA;
                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-US20327.
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 (first entry)
                                                                                                                                                                                                            97.1%;
95.5%;
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                                                                 22 AA
                                                                                                                                                                                                             Score 102; DB 20
Pred. No. 6e-08;
                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                               DB 20;
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                                                                                                                                                                                                                               Length 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia;
                                        Lecithin: cholesterol acyltransferase activation exhibiting peptide #169.
                                                                                                                 14-JUL-1999
                                                                                                                                                            AAY19356;
                                                                                                                                                                                           AAY19356 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an compliance, are used to treat or prevent diseases associated with disease, are used to treat or prevent diseases associated with disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that e.g. atherosclerotic streaks, and to raise antibodies. AAY19188 to avhibition constitute the constitution of the constitut
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 124; 280pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide agonists of apolipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-277035/23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (METZ/)
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                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemla; metabolic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I;
                                                                                                                                                                                                                                                                                                                    ۲
                                                                                                                                                                                                                                                                                                                                                1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia;
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                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MET2 G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cornut I,
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95.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                            Score 102; DB 2
Pred. No. 6e-08;
                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            DB 20; Length 22;
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                                                                                                                                                                                                                                                                                                              Matches
              Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin; cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I;
 septic shock
high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                                                                                                                    09-JUL-1999
                                                                                                                                                                      AAY19112 standard; Peptide;
                                                                      Lecithin:cholesterol acyltransferase activation exhibiting peptide #179.
                                                                                                                                                   AAY19112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that
                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                              is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY19188 to AAY19441 represent lecithin:cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (apoA I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an agonist (A) of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 125; 280pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide agonists of apolipoprotein A-I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-277035/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Buttner K, Cornut I,
                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                              l pvlelfenlleklldalqkklk 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
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                                                                                                                                                                                                                                                                            1 PVLELFENLLERLLDALQKKLK 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DUFOURCQ J
                                                                                                                                                                                                                                                                                                                                                                                22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                     core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                               (first entry)
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95.5%;
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                                                                                                                                                                         22 AA
                                                                                                                                                                                                                                                                                                                 Score 102; DB 2
Pred. No. 6e-08;
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Best Local .
           WO9916458-A1
                                                                                      human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I;
                                 Homo sapiens
                                                  Synthetic.
                                                                           septic shock
                                                                                     high density lipoprotein;
                                                                                                                  Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia
                                                                                                                                            Lecithin: cholesterol acyltransferase activation exhibiting peptide #188
                                                                                                                                                                            09-JUL-1999 (first entry)
                                                                                                                                                                                                          AAY19121;
                                                                                                                                                                                                                            AAY19121 standard; Peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 apoA-I deficiency: hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to AAY19187 represent lecithin; cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                         exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein)
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 116; 254pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide agonists of apolipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-277034/23
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                                                                                                                                                                                                                                                                                                                        1 PVLELFENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                               21;
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SEKUL R.
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                                                                                                                                                                                                                                                                                                                                                                                                                     22 AA;
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95.5%;
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                                                                               hypertriglyceridemia; metabolic syndrome
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Pred. No. 6e-08;
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AAY19102
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Best Local
                                                                                                                                                        Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosclerosis; restenosis; HDL; apoA-I; high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
29-SEP-1997;
                            28-SEP-1998;
                                                        08-APR-1999
                                                                               W09916458-A1
                                                                                                                                                     septic shock
                                                                                                          Homo sapiens
                                                                                                                                                                                                                            Lecithin: cholesterol acyltransferase activation exhibiting peptide #169.
                                                                                                                                                                                                                                                              09-JUL-1999 (first entry)
                                                                                                                                                                                                                                                                                             AAY19102;
                                                                                                                                                                                                                                                                                                                     AAY19102 standard; Peptide; 22 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treating septic shock. When labeled, (A) can also be used diagnostically to measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to AAY19187 represent lecithin cholesterol acyltransferase (LCAT) activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular disease, atherosclerosis, restenosis, HDL (high density lipoprotein) or apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 18; Page 117; 254pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide agonists of apolipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-277034/23.
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(DASS/) DASSEUX J.
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95.5%;
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Pred. No. 6e-08;
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AAY19083
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Best Local :
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(CORN/) CORNUT I.
(DASS/) DASSEUX J.
(METZ/) METZ G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Apolipoprotein A-I; agonist; dyslipidemic disorder; dyslipidemia; human; lecithin:cholesterol acyltransferase; LCAT; hypercholesterolaemia; cardiovascular disease; atherosolerosis; restenosis; HDL; apoA-I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lecithin:cholesterol acyltransferase activation exhibiting peptide #150.
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                                                                                                                                                                                                                                                                 28-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   septic shock.
                                                                                                                                                                                                      29-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            high density lipoprotein; hypertriglyceridemia; metabolic syndrome;
                           (SEKU/)
                                                                                   (DASS/) DASSEUX J.
                                                                                                               (CORN/)
                                                                                                                                           (BUTT/) BUTTNER K.
                                                           (METZ/) METZ G.
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                           SEKUL R.
                                                                                                                  CORNUT I
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Pred. No. 6e-08;
1; Mismatches
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Job time:

Search completed: November 20, 2001,

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                                                                                                                                                                                            The present invention describes an agonist (A) of apolipoprotein A-I (apoA-I) which is a 15-29 residue peptide, or analog, that forms an amphipathic alpha-helix in presence of lipids. (A), and their lipid complexes, are used to treat or prevent diseases associated with dyslipidemia, specifically hypercholesterolaemia, cardiovascular apoA-I deficiency; hypertriglyceridemia and metabolic syndrome, also for treating septic shock. When labeled, (A) can also be used diagnostically treating septic shock. When labeled, (A) can also be used diagnostically tro measure serum levels of HDL, in particular the HDL subpopulation that is involved in retrograde cholesterol transport, also to image HDL at e.g. atherosclerotic streaks, and to raise antibodies. AAY18934 to AAY19187 represent lecithin:cholesterol acyltransferase (LCAT) activity exhibiting core peptides, which are apoA-I agonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 18; Page 115; 254pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide agonists of apolipoprotein A-I
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                                                                                                                                                                 Sequence
1 PVLELFENLLERLLDALQKKLK 22
                                                                            21;
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                                                                          Conservative
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                                                                                                Score 102; DB 2
Pred. No. 6e-08;
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Result
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Copyright (c) 1993 - 2000 Compugen Ltd.
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S02788
A23654
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Best Local Similarity
Whiches 7; Conserve
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A:Gene: HP0560; jhp0507
                                                                                                                                                                                                                                                                                                                                       A:Title: Genomic sequence comparison of two unrelated isolates of the human gastric p A;Reference number: A71800; MUID:99120557
A;Recession: A71923
A;Molecule type: DNA
A;Residues: 1-26 <ARN>
A;Residues: 1-26 <ARN>
A;Cross:references: GB:AE001484; GB:AE001439; NID:g4155043; PIDN:AAD06083.1; PID:g415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A:Cross-references: GB:AE000570; GB:AE000511; NID:g2313672; PIDN:AAD07633.1; PID:g231A; Experimental source: strain 26695
R;Alm, R.A.; Ling, L.S.L.; Moir, D.T.; King, B.L.; Brown, E.D.; Doig, P.C.; Smith, D. Ives, C.; Gibson, R.; Merberg, D.; Mills, S.D.; Jiang, Q.; Taylor, D.E.; Vovis, G.F. Nature 397, 176-180, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Species: Helicobacter pylori
A;Variety: Strains J99, 26695
C;Jate: 09-Aug-1997 #sequence_revision 09-Aug-1997 #text_change 08-Oct-1999
C;Accession: H64589; A71923
R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKe son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey, Nature 388, 539-547, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status: nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Residues: 1-26 <TOM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A:Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A:Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A:Reference number: A64520; MUID:97394467
A:Reference number: A64520; MUID:97394467
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H64589
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15 LLYRILDVLEQK 26
                                                    9 LLERLLDALQKK 20
                                                                                                                     Conservative
                                                                                                                                          33.3%;
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                                                                                                                                          Score 35;
Pred. No.
                                                                                                         Pred. No. 97;
3; Mismatches
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                                                                                                                                                                      Length 26;
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N5.N10-methenyltetrahydromethanopterin cyclohydrolase - Methanosarcina barkeri (fragm C:Species: Methanosarcina barkeri C:Date: 19-Nov-1993 #sequence_revision 18-Nov-1994 #text_change 18-Nov-1994 C:Accession: B48368 R:Klein, A.R.; Breitung, J.; Linder, D.; Stetter, K.O.; Thauer, R.K. Arch. Microbiol. 159, 213-219, 1993 A:Title: N5.N10-methenyltetrahydromethanopterin cyclohydrolase from the extremely the

RESULT B48368

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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-24 <KLE>
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C; Geneti
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C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 31-Mar-2001
C;Accession: B85840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Genome sequence of enterchemorrhagic Escherichia coli 0157:H7 A;Reference number: A85480; MUID:21074935; PMID:11206551 A;Accession: B85840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B85840
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A;Molecule type: DNA
A;Residues: 1-29 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession: C85840
R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; Mayhewiller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; Apodaca, Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypothetical protein Z3290 [imported] - Escherichia coli (strain O157:H7) C;Species: Escherichia coli (c;Species: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 31-Mar-2001 C;Date: 16-Feb-2001 #sequence_revision 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:AE005174; NID:g12516340; PIDN:AAG57182.1; GSPDB:GN00145; UWGP:23; A;Experimental source: strain 0157:H7, substrain EDL933
                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Genome sequence of enterohemorrhagic Escherichia coli O157:H7 A;Reference number: A85480; MUID:21074935; PMID:11206551 A;Accession: C85840
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                                                                                                                                                                                                                              A;Cross-references: GB:AE005174; NID:g12516341; PIDN:AAG57183.1; GSPDB:GN00145; UWGP:Z3:A;Experimental source: strain 0157:H7, substrain EDL933
                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-29 <STO>
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58.8%;
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Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 31.5; DB 2;
Pred. No. 3.4e+02;
                                   Pred. No. 3.4e+02;
                                                               Score 31.5;
Mismatches
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                                                                   DB 2; Length 29;
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A; Molecule type: protein
A; Note: the residues: 1-2,'xx',5-9,'x',11-13,'x',15-21 <HAA>
A; Note: the residues designated 'X' were determined to be gamma-carboxy glutamic acid
C; Comment: This venom peptide induces sleep in young mice and is an NMDA antagonist i
C; Superfamily: conantokin
C; Keywords: amidated carboxyl end; carboxyglutamic acid; neurotoxin; venom
F; 31,4,10,14/Modified site: gamma-carboxyglutamic acid (Glu) #status experimental
F; 21/Modified site: amidated carboxyl end (Ala) #status experimental
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A;Title: Conantokin-T. A gamma-carboxyglutamate containing peptide with N-methyl-D-as A;Reference number: A35225; MUID:90202866
A;Accession: A35225
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C:Accession: A35225
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C; Species: Conus tulipa (tulip cone)
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A; Molecule type: protein
A; Residues: 1-30 < KRW>
C; Superfamily: corticotropin-lipotropin
C; Superfamily: corticotropin-lipotropin
C; Keywords: acetylated amino end; opioid peptide
F;1-5/Product: Met-enkephalin *status predicted <MEN>
F;1/Modified site: acetylated amino end (Tyr) *status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biochem. Biophys. Res. Commun. 92, 1278-1288, 1980 A;Title: Occurrence of two different endorphins in the salmon A;Reference number: A01469; MUID:80174902 A;Accession: A01469
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   A; Variety: strain HST
C; Date: 21-Jan-2000 #:
C; Accession: T43971
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                                                                                                      C; Species: human herpesvirus 6
                                                                                                                                          CC-chemokine receptor [imported] - human herpesvirus 6 (strain HST)
                                                                                                                                                                             T43971
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Best Local :
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          21-Jan-2000 #sequence_revision 21-Jan-2000 #text_change 21-Ju1-2000
sion: T43971
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pred. No. 2.8e+02;
4; Mismatches 6;
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Pred. No. 4.2e+02;
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RESULT 9
PLO027
M protein pepM19 - Streptococcus sp. (fragment)
C;Species: Streptococcus sp.
C;Species: Streptococcus sp.
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C;Keywords:
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                                                                                                                                                                                                                                                      A; Reference number: PL0027; MUID:88258373
                                                                                                                                                                                                                                                                J. Exp. Med. 167, 1849-1859, 1988
A;Title: Protective and heart-crossreactive epitopes located within the NH2 terminus
                                                                                                                                                                                                                                                                                                     R;Bronze, M.S.; Beachey, E.H.; Dale, J.B
J. Exp. Med. 167, 1849-1859, 1988
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A:Residues: 1-25 <WAD>
C:Comment: This hydrogen peroxide-scavenging peroxidase plays a
C:Keywords: oxidoreductase
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R;Wadda, N.; Kinoshita, S.; Matsuo, M.; Amako, K.; Miyake, C.; Asada, K.
Biochem. Biophys. Res. Commun. 242, 256-261, 1998
A;Title: Purification and molecular properties of ascorbate peroxidase from bovine eye.
A;Reference number: PC4445; MUID:98113338
A;Accession: PC4445
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A;Title: Comparison of the complete DNA sequences of human herpesvirus 6 variants A and A;Reference number: Z22732; MUID:99412319
A;Accession: T43971
A;Scatus: preliminary; translated from GB/EMBL/DDBJ
A;Rolecule type: DNA
A;Residues: 1-24 <ISE>
A;Cross:references: EMBL:AB021506; NID:9495977; PIDN:BAA78232.1; PID:94995999
A;Experimental source: Strain HST; pop. variant B
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C:Species: Bos primigenius taurus (cattle)
C:Date: 28-Mar_1998 #sequence_revision 08-May-1998 #text_change 07-May-1999
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Best Local :
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Best Local Similarity
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EDKLKKIIDDLDAK 22
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71.4%;
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50.0%;
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                                                                                     Score 27; DB 2;
Pred. No. 1.3e+03;
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Pred. No.
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Pred. No. 1.2e+03;
3; Mismatches 3;
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                                                                                                     Length 26;
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       Matches
                     Best Local Similarity
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     Conservative
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A;Molecule type: protein
A;Residues: 1-24 <LAM>
C;Comment: Limulus polyphemus hemocyanin is an association of eight different subunit
                                                                                                                 A;Title: Quaternary structure of Limulus polyphemus hemocyanin A;Reference number: A90478
                                                                                                                                                                             C;Accession: H20554
R;Lamy, J.; Lamy, J.; Sizare
Biochemistry 22, 5573-5583,
                                                                                                                                                                                                                                           hemocyanin subunit VI - Atlantic horseshoe crab (fragment)
C:Species: Limulus polyphemus (Atlantic horseshoe crab)
C:Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 08-Jul-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-24 <LAM>
C;Comment: Limulus polyphemus hemocyanin is an association of eight different subunit
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A;Accession: A20554
A;Accessionet type: protein
A;Residues: 1-24 <LAM>
C;Comment: Limulus polyphemus hemocyanin is an association of eight different subunit
C;Superfamily: hemocyanin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Limulus polyphemus (Atlantic horseshoe crab)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 08-Jul-1994
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C;Species: Limulus polyphemus (Atlantic horseshoe crab)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 12-Apr-1995
C;Accession: A20554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hemocyanin subunit V - Atlantic horseshoe crab (fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Title: Quaternary structure of Limulus polyphemus hemocyanin
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62.5%;
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Pred. No.
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Pred No. 1.7e+03;
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24.8%; 50.0%;

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Score 26; DB 2; Pred. No. 1.7e+03 4; Mismatches

Length 24

Indels

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Gaps

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DP 6A

13 LLDALQKKLK 22 :| ||:|:|: 1 VLGALEKQLR 10

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A; Molecule type: protein
A; Residues: 1-26 <FED>
-C; Superfamily: gastric intrinsic factor
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S02788
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Biochim. Biophys. Acta 1292, 113-119, 1996
A;Title: Transcobalamin from cow milk: isolation and physico-chemical properties. A;Reference number: S62672; MUID:96139334
A;Accession: S62672
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C;Species: Bos primigenius taurus (cattle)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 02-Jul-1998
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ceptor-related protein.
A; Reference number: A23691; MUID:91093092
A; Accession: A23691
A; Status: preliminary
A; Molecule type: protein
A; Residues: 1-28 <WEI'>
C; Superfamily: apolipoprotein A-I
C; Keywords: lipid binding; lipoprotein
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A;Residues: 1-17;18-26 <GRA>
C;Keywords: acyltransferase; coenzyme A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Grant, D.M.; Lottspeich, F.; Meyer, U.A. FEBS Lett. 244, 203-207, 199 A;Title: Evidence for two closely related isozymes of arylamine N-acetyltransferase in A;Reference number: S02788; MUID:89171260
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                                                                                                                                                                                                                              R;Weisgraber, K.H.; Mahley, R.W.; Kowal, R.C.; Herz, J.; Goldstein, J.L.; Brown, M.S. J. Biol. Chem. 265, 22453-22459, 1990
A;Title: Apolipoprotein C-I modulates the interaction of apolipoprotein E with beta-migrantic.
                                                                                                                                                                                                                                                                                                                  apolipoprotein C-I - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 04-Oct-1991 #sequence_revision 04-Oct-1991 #text_change 31-Dec-1993
C;Accession: A23691
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Best Local Similarity 40.0%;
Matches 6; Conservative
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Best Local !
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Pred. No. 1.8e+03;
4; Mismatches 4
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Query Match
Best Local Similarity 30.0%; Pred. No. 2e+03;
Matches 6; Conservative 6; Mismatches 8; Indels 0; Gaps 0;

Qy 3 LELFENLLERLLDALQKKLK 22
| | | : | : | : | : |
Db 7 LELIPDKLKEFGNTLEEKAR 26
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Search completed: November 20, 2001, 09:43:45 Job time: 159 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
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Match
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Copyright (c) 1993 - 2000 Com
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atch		STRUCTURE BY NM MEDLINE=9715300 Skjaerbaek N., Zbetermination "Determination conantokin T by J. Biol. Chem:-FUNCTION: I N.METHYL-D- CENTRAL NER PIR: A35225; A3 PDB: 10NT; 04-S Venom; Vitamin 3D-structure.	E SE	SEQUENCE. SEQUENCE. TISSUE=Venom: MEDLINE=902008: Haack J.A., R: Olivera B.M.; "Conantokin-T "Conantokin-T J. Biol. Chem	(N)	1 FU F_CONTU	22 21.5 21.5 21 21 21 21 21 21 21 21 21 21 21 21 21
Similarity 7; Conserva	3 4 10 14 21 2 AA;	TURE BY NMR. NE=97153002; PubMed= rbaek N., Nielsen K. rmination of the sol tokin-T by CD and NM col. Chem. 272:2291-2 UNCTION: INDUCES SLE -METHYL-D-ASPARTATE ENTRAL NERVOUS SYSTE A35225, A35225. 10NT; 04-SEP-97. ; Vitamin K; Gamma-c ructure.	· (+	; 2866; Rivie ; ; T. A ; spart; m. 26;	Sh.	STAN	211.0 221.0 20.0 20.0 20.0 20.0 20.0 20.
29.5% 41.2% tive	3 4 10 14 21 2509 MW;	IR. Nielsen K.J., Le of the solution CD and NMR spec 272:2291-2299(19 NDUCES SLEEP-LIK ASPARTATE (NNDA) VOUS SYSTEM NEUR 5225. K; Gamma-carboxy K; Gamma-carboxy	bMed=9 , Zhu ructur absen (1997)	PubMed=2 er J.E., F gamma-car gate antag	Las Las Intin Intin	TANDARD;	27 1 28 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
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21-JUL-1986 (Rel. 0
01-NOV-1990 (Rel. 1
BETA-ENDORPHIN II
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID-8018;
                                                                                                                                                                                                                                                                                                                                 NUO6_SOLTU STANDARD; PRT; 19 AA. P80729; P80729; O1-NOV-1997 (Rel. 35, Created) O1-NOV-1997 (Rel. 35, Last sequence update) O1-NOV-1997 (Rel. 35, Last annotation update) NADH-UBIQUINONE OXIDOREDUCTASE 11 KDA SUBUNIT (EC 1.6.5.3) (EC 1.6.99.3) (COMPLEX I-11KD) (CI-11KD) (FRAGMENT).
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                                                                                                                                                                          Herz U., Grohmann L.;
Submitted (DEC-1996) to the SWISS-PROT data bank.
-i- FUNCTION: TRANSFER OF ELECTRONS FROM NADH TO THE RESPIRATORY
CHAIN. THE IMMEDIATE ELECTRON ACCEPTOR FOR THE ENZYME IS BELIEVED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   END2_ONCKE
P01205;
                                                                                Oxidoreductase; NAD; Ubiquinone; Mitochondrion. NON_TER 19 19
                                                                                                                                                                                                                                                                                   Solanum tuberosum (Potato).

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;

Magnoliophyta; eudicotyledons; core eudicots; Asteridae; euasterids

Solanales; Solanaceae; Solanum.
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                                                                                                                    TO BE UBIQUINONE.

CATALYTIC ACTIVITY: NADH + UBIQUINONE = NAD(+) + UBIQUINOL.
SUBUNIT: COMPLEX I IS COMPOSED OF ABOUT 30 DIFFERENT SUBUNIT:
SUBCELLULAR LOCATION: MATRIX SIDE OF THE MITOCHONDRIAL INNER
                                                                                                          MEMBRANE.
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                                                                  2310 MW;
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35.7%;
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Score 29; DB Pred. No. 2.4e 5; Mismatches
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Eur. J. Biochem. 219:145-154(1994).

-i- FUNCTION: POSSESSES A POTENT ANTIMICROBIAL ACTIVITY AGAINST
BACTERIA FUNGI AND PROTOZOA. PROBABLY ACTS BY DISTURBING MEMBRANE
FUNCTIONS WITH ITS AMPHIPATIC STRUCTURE.

-i- SUBCELLULAR LOCATION: SECRETED.
-i- TISSUE SPECIFICITY: SKIN.
Antibiotic; Fungicide; Multigene family; Amphibian skin.
SEQUENCE 27 AA; 2779 MW; 43C94D2DC19721A8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P56920;
30-MAY 2000 (Rel. 39, Created)
01-0CT-2000 (Rel. 40, Last seq
01-0CT-2000 (Rel. 40, Last ann
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Ol-FEB-1994 (Rel. 28, Created)
Ol-FEB-1994 (Rel. 28, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
DERMASEPTIN 4 (DS IV).
Phyllomedusa sauvagei (Sauvage's leaf frog).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rana temporaria (European common frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea;
NCBI_TaxID=8407;
                                                                                                                       "Isolation and structure of novel defensive peptides from frog skin."; Eur. J. Biochem. 219:145-154(1994).
                                                                                                                                                      Mor A., Nicolas P.;
                                                                                                                                                                    MEDLINE=94139686; PubMed=8306981;
                                                                                                                                                                                  TISSUE-Skin;
                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                                                                                            Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            VLPIIGNLLNSLL 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biochem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antimicrobial peptides from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mignogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     242:788-792(1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26.7%;
53.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Canofeni S., Miele R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence update) annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 28;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AMIDATION.
26505DFA79A92448 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                        update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 European red frog Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 13
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APCL_RABIT
ID APCLE,
AC P33047
DT 01-OCT
DT 15-DEC
DE APOLIP
GN APOCLIO
OS Orycto
OC Eukary
OC Mammal
OX MCBI_T
RN [1]
RN [1]
RN MEDLIN
RA Weisgr
RA Weisgr
RA Weisgr
RA Trecept
RT with b
RT inhibi
RT recept
CC -i- FU
CC -i- FU
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CC -i- SI
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Y425_B
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Best Local Similarity
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Best Local
                                                                                                                                                                                                                                                                                           15-DEC-1998
15-DEC-1998
15-DEC-1998
     SEQUENCE FROM N.A.
STRAIN=ATCC 35210 / B31;
MEDLINE=98065943; PubMed=9403685;
Fraser C.M., Casjens S., Huang W.M.,
Lathigra R., White O., Ketchum K.A.,
                                                                                                                                                                                                                                                                                                                                                                      051386;
                                                                                                                                                                                                                                                                                                                                                                                                                        BORBU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Apolipoprotein C-I modulates the interaction of apolipoprotein E with beta-migrating very low density lipoproteins (beta-VLDL) and inhibits binding of beta-VLDL to low density lipoprotein receptor-related protein.";

J. Biol. Chem. 265:22453-22459(1990).

-i- FUNCTION: APPEARS TO MODULATE THE INTERACTION OF APOE WITH BETA-MIGRATING VLDL AND INHIBIT BINDING OF BETA-VLDL TO THE LDL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-1993 (Rel. 27, Created)
01-OCT-1993 (Rel. 27, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
                                                                                                                                                                                 Borrelia burgdorferi (Lyme disease spirochete)
Bacteria; Spirochaetales; Spirochaetaceae; Bor
                                                                                                                                                                                                                                                              HYPOTHETICAL PROTEIN BB0425.
                                                                                                                                                            NCBI_TaxID=139;
                                                                                                                                                                                                                                            BB0425
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plasma; Lipid transport; VLDL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIR; A23691; A23691.
HSSP; P02654; 10PP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brown M.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Weisgraber K.H., Mahley R.W., Kowal R.C., Herz J., Goldstein J.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=91093092; PubMed=2266137;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APOLIPOPROTEIN C-I (APO-CI) (FRAGMENT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APC1_RABIT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LFENLLERLLDALQKKL 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28 28
28 AA; 3165 MW;
                                                                                                                                                                                                                                                                                 (Rel. 37, Created)
(Rel. 37, Last sequence up
(Rel. 37, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24.8%;
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41.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 26; DB 1; pred. No. 9.3e+02; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2903C027F676541C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 28; DB 1;
Pred. No. 4.8e+02;
4; Mismatches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                           update)
Sutton G.G., Clayton R.A., Dodson R., Hickey E.K., Gwinn
                                                                                                                                                                                                                                                                                   update)
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                                                                                                                                                                                    Borrelia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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TEMK_RA
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""" 6; Conserv
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                                                                         Query Match
Best Local (
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01-0CT-2000 (Rel. 40, Last sequence update)
01-0CT-2000 (Rel. 40, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                   Rana temporaria (European common frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dougherty B., Tomb J.-F., Fleischmann R.D., Richardson D., Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M. van Vugt R., Palmer N., Adams M.D., Gocayne J.D., Weidman J., Utterback T., Watthey L., McDonald L., Artiach P., Bowman C., Garland S., Fujii C., Cotton M.D., Horst K., Roberts K., Hatch B., Smith H.O., Venter J.C.;
                                                                                                                                        SEQUENCE 10 AA; 1123 MW;
                                                                                                                                                     Amphibian skin; Antibiotic; Amidation; Multigene family MOD_RES 10 10 AMIDATION.
                                                                                                                                                                                                                     -!- SUBCELLULAR LOCATION: SECRETED
                                                                                                                                                                                                                                                                              "Temporins, antimicrobial peptides from the European red frog Rana temporaria.";
                                                                                                                                                                                                                                                                                                                 Barra D.,
                                                                                                                                                                                                                                                                                                                           Simmaco M., Mignogna
                                                                                                                                                                                                                                                                                                                                             MEDLINE=97175050; PubMed=9022710;
                                                                                                                                                                                                                                                                                                                                                               TISSUE=Skin;
                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=8407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P56923;
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hypothetical protein. SEQUENCE 30 AA; 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                     -! - SIMILARITY: BELONGS TO
                                                                                                                                                                                                                                                 -!- FUNCTION: HAS ANTIBACTERIAL ACTIVITY AGAINST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TEMK_RANTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Genomic sequence of a Lyme disease spirochaete, Borrelia
                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 QNLLIFLNKKIK 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 ERLLDALQKKLK 22
                             Ç
1 LLPNLLKSLL 10
                                                                                                                                                                                      FAMILY.
                                                                                                                                                                                                                                    BACTERIA.
                           LFENLLERLL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BB0425; -
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                                                                                                                                                                                                                                                                 Biochem. 242:788-792(1996)
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
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                                                                         23.8%;
                                                                                                                                                                                                                                                                                                                           G., Canofeni S., Miele R., Mangoni M.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24.8%;
50.0%;
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                                                         Pred. No. 4.3e+02
l; Mismatches
                                                                                                                                      390549B337272457 CRC64;
                                                                                        Score 25;
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Pred. No. 1e+03;
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                                                         Indels
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NIFD
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Best Local
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P33177;
01-OCT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BIOCHIM. BIOPHYS. ACTA 1171:337-340(1993).

BIOCHIM. BIOPHYS. ACTA 1171:337-340(1993).

FUNCTION: THE KEY ENZYMATIC REACTIONS IN NITROGEN FIXATION ARE CATALYZED BY THE NITROGENASE COMPLEX, WHICH HAS 2 COMPONENTS: THOO PROTEIN.

IRON PROTEIN AND THE MOLYBDENUM-IRON PROTEIN.

ICATALYTIC ACTIVITY: 8 REDUCED FERREDOXIN + 8 H(+) + N(2) + 16 AV + 16 AV + 16 ORTHOPHOSPHATE.

ICATALYTIC ACTIVITY: 8 REDUCED FERREDOXIN + 2 NH(3) + 16 ADP + 16 ORTHOPHOSPHATE.

ICATALYTIC ACTIVITY: BELONGS TO THE NIFD/NIFK/NIFE/NIFN FAMILY.

ICATALYTIC ACTIVITY: BELONGS TO THE NIFD/NIFK/NIFE/NIFN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-0CT-1993 (Rel. 27, Created)
01-0CT-1993 (Rel. 27, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
NITROGENASE MOLYBDENUM-IRON PROTEIN ALPHA CHAIN (EC 1.18.6.1)
                                                                                                                                                                                                                                  CT21_LITCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Murphy S.T., Jackman D.M., Mulligan M.E.; "Cloning and nucleotide sequence of the gene reductase (nifH) from the heterocyst-forming
                                                                                                                                            CT21_LITCI STANDARD; PRT; 26 AA.
P81847; P81848; P81849; P81850;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
CITROPIN 2.1.3 [CONTAINS: CITROPIN 2.1.2; CITROPIN 2.1.1; CITROPIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. MEDLINE=93144353; PubMed=8424961;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anabaena sp. (strain L31).
Bacteria; Cyanobacteria; Nostocales; Nostocaceae; Anabaena
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oxidoreductase; Nitrogen fixation; Molybdenum; Iron-sulfur.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS00090; NITROGENASE_1_2; PARTIAL. PROSITE; PS00699; NITROGENASE_1_1; PARTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; L04499; AAA22015.1; -. PIR; S28191; S28191.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=29412;
         SEQUENCE
                                      NCBI_TaxID=94770;
                                                                        Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
                                                                                             Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000318; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NITROGENASE COMPONENT I) (DINITROGENASE) (FRAGMENT).
                                                                                                           Litoria citropa (Australian blue mountains tree frog).
                                                                                                                                                                                                                                                                                                                       7 KNLVQENKELIQEVLK 22
                                                                                                                                                                                                                                                                                                                                                          ENLLERLLDALQKKLK 22
                                                                                                                                                                                                                                                                                                                                                                                           Similarity
5; Conserv
                                                                                             Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23 AA; 2666 MW;
                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                             Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                           23.8%;
                                                                                                                                                                                                                                                                                                                                                                                             6;
                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; DB 1;
Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9F2C70302E167806 CRC64;
                                                                                             Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23
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                                                                                                                                                                                                                                                                                                                                                                                                                            Length 23;
                                                                                              Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anabaena
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AC DT DT DT

TEMD_RANTE P56919;

STANDARD;

PRT;

13

30-MAY-2000

01-OCT-2000 01-OCT-2000 TEMPORIN D.

(Rel. (Rel. (Rel.

39, Created)
40, Last sequence update)
40, Last annotation update)

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TEMD_RANTE
                                                    B
            RESULT 12
                                                                                                      Query Match
Best Local S
Matches 6
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Best Local Similarity
Matches 5; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wegener K.L., Wabnitz P.A., Carver J.A., Bowie J.H., Chia B.C.S., Wallace J.C., Tyler M.J.;
"Host defence peptides from the skin glands of the Australian blue mountains tree-frog Litoria citropa. Solution structure of the antibacterial peptide citropin 1.1.";
Eur. J. Biochem. 265:627-637(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                               p56918;
30-MAY-2000 (Rel. 39, Created)
01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                  TEMPORIN C.
Rana temporaria (European common frog).
Runa temporaria (European common frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Neobatrachia; Ranoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EUR. J. BIOCHGHI. 2017Y: DORSAL AND SUBMENTAL
-1- TISSUE SPECIFICITY: DORSAL AND SUBMENTAL
-1- TISSUE SPECIFICITY: DORSAL AND SUBMENTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=99435977; PubMed=10504394;
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                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                        Amphibian skin; Antibiotic; Amidation; Multigene family MOD_RES 13 13 AMIDATION.
                                                                                                                                                                                                                                                                                          "Temporins, antimicrobial peptides from the temporaria.";
                                                                                                                                                                                                                                                                                                                         Barra D.;
                                                                                                                                                                                                                                                                                                                                     Simmaco M.,
                                                                                                                                                                                                                                                                                                                                                   MEDLINE=97175050; PubMed=9022710;
                                                                                                                                                                                                                                                                                                                                                               TISSUE=Skin;
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=8407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TEMC_RANTE
                                                                                                                                                                                                                                                                 -!- FUNCTION: HAS ANTIBACTERIAL ACTIVITY AGAINST GRAM-POSITIVE
                                                                                                                                                                                                                           -i- SIMILARITY: BELONGS TO THE BREVININ/ESCULENTIN/GAEGURIN/RUGOSIN
                                                                                                      Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 LVDVLKPKLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13 LLDALOKKLK 22
                                                     1 LLPILGNLLNGLL
                                                                              2 VLELFENLLERLL 14
                                                                                                                                                                                                                   FAMILY.
                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: SECRETED
                                                                                                                                                                                                                                                         BACTERIA
                                                                                                                                                                                                                                                                                   J. Biochem. 242:788-792(1996)
                                                                                                                                                                          13 AA; 1363 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26
                                                                                                                                                                                                                                                                                                                                   Mignogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23
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26
2519 MW;
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                                                                                                                                                                                                                                                                                                                                      G., Canofeni S., Miele R., Mangoni M.L.,
                                                                                                                      22.9%;
46.2%;
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CITROPIN 2.1.1.
CITROPIN 2.1.2.
CITROPIN 2.1.3.
CITROPIN 2.1.3.
7490C28EB2E4899A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 25; DB 1; LC Pred. No. 1.2e+03; Wismatches 2;
                                                                                                                      Score 24; I
Pred. No. 7
                                                                                                                                                                           2201403A655B2448 CRC64
                                                                                                                                                                                      AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         update)
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                                                                                                                                     DB 1;
                                                                                                          .8e+02;
les 5;
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                                                                                                                                  Length 13;
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                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                        Ranidae;
                                                                                                                                                                                                                                                                                                              red frog Rana
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KT39_P
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Best Local .
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                                                             01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOX in.
Ovis aries (Sheep)
                             SCT
                                                   SECRETIN
                                                                                                                                           SECR_SHEEP
P31299;
                                                                                                                                                                                            SHEEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pichia kluyveri (Yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Pichia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KT39_PICKL STANDARD; PRT; 27 AA. P80326; PR0326; PRO (Control of the control of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=ATCC 66811 / GBS 7145;
Martin C., Leis W., Radler F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PICKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amphibian skin; Amidation; Multigene family. MOD_RES 13 13 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Simmaco M., Mignogna G., Canofeni S., Miele R., Mangoni M.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eur. J. Biochem. 242:788-792(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          temporaria."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Barra D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=97175050; PubMed=9022710;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rana temporaria (European common frog).
Eukaryota: Metazoa: Chordata: Craniata: Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Temporins, antimicrobial peptides from the European red frog Rana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-Skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE, AND SYNTHESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=8407;
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                                                                                                                                                                                                                                                                                 16 ENLAQQFVD 24
                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                7 ENLLERLLD 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: SECRETED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBCELLULAR LOCATION: SECRETED.
SIMILARITY: BELONGS TO THE BREVININ/ESCULENTIN/GAEGURIN/RUGOSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LLPIVGNLLNSLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27 AA;
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27
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                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3084 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Radler F.;
                                                                                                                                                                                                                                                                                                                                                                                          22.98;
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46.2%;
                                                                                                                                                                                                                                                                                                                                                                                          Score 24;
Pred. No. 1
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Pred. No.
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3EF35DFA655B2448 CRC64;
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                                                                 update)
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1.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                              Length 27;
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                      FUNCTION: PROBABLE BIFUNCTIONAL ENZYME SHOWING MINIMAL GLUTATHIONE-CONJUGATING ACTIVITY WITH ETHACRYNIC ACID AND 7-CHLORO-4-NITROBENZ-2-OXA-1, 3-DIAZOLE AND MALEYLACETOACETATE ISOMERASE ACTIVITY. HAS ALSO LOW GLUTATHIONE PEROXIDASE ACTIVITY WITH T-BUTYL AND CUMENE HYDROPEROXIDES (BY SIMILARITY). IS ABLE TO CATALYZE THE GLUTATHIONE DEPENDENT OXYGENATION OF DICHLOROACETIC ACID TO GLYOXYLIC ACID.

-!- CATALYTIC ACTIVITY: 4-MALEYLACETOACETATE = 4-FUMARYLACETOACETATE.
-!- COFACTOR: THE MAAI ACTIVITY REQUIRES GLUTATHIONE (BY SIMILARITY).
-!- PATHWAY: CATABOLISM OF TYROSINE; FOURTH STEP, CATABOLISM OF TROSINE;
                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-2000 (Rel. 40, Created)
01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
MALEYLACETOACETATE ISOMERASE (EC 5.2.1.2) (MAAI) (GLUTATHIONE TRANSFERASE ZETA 1) (EC 2.5.1.18) (GSTZ1-1) (FRAGMENT).
                                                                                                                                                                                               "Glutathione transferase zeta catalyses the oxygenation carcinogen dichloroacetic acid to glyoxylic acid.", Biochem. J. 331:371-374(1998).

-I- FUNCTION: PROBABLE BIFUNCTIONAL ENZYME SHOWING MINING.
                                                                                                                                                                                                                                                         MEDLINE=98198370; PubMed=9531472; Tong Z., Board P.G., Anders M.W.;
                                                                                                                                                                                                                                                                                          SEQUENCE, AND CHARACTERIZATION.
STRAIN=FISCHER 344; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID-10116;
                                                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P57113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00260; GLUCAGON; 1.
Glucagon family; Hormone; Amidation.
MOD_RES 27 27 AMIDATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00123; hormone2; 1.
PRINTS; PR00275; GLUCAGON.
PROSITE; PS00260; GLUCAGON.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PIR; C60072; SESH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide, peptide histidine isoleucinamide and secretin from the ovine small intestine.";
Regul. Pept. 32:169-179(1991).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
Mammalia; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000532; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Purification and amino acid sequence of vasoactive intestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bounjoua Y., Vandermeers A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=91239834; PubMed-2034821;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Small intestine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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     SUBUNIT: HOMODIMER
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                  PHENYLALANINE; FIFTH STEP
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SIMILARITY).
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Pred. No.
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CC -!- SUBCELLULAR LOCATION: CYTOSOPLASMIC.
CC -!- PTM: THE N-TERMINUS IS BLOCKED (PROBABLE).
CC -!- SIMILARITY: BELONGS TO THE GST SUPERFAMILY. ZETA FAMILY.
CC |-- SIMILARITY: BELONGS TO THE GST SUPERFAMILY. ZETA FAMILY.
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Title:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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"Genomic sequence comparison of two unrelated isolates of the human gastric pathogen Helicobacter pylori."; Nature 397:176-180(1999).	Gibson R., Merberg D., Mills S.D., Jiang Q., Taylor D.E., Vovis G.F., Trust T.J.;	Alm K.A., Ling L. S.L., Moir D.T., King B.L., Brown E.D., Doig P.C., Smith D.R., Noonan B., Guild B.C., deJonge B.L., Carmel G., Timming B., Carmel G., Timming B., Carmel G.,	MEDLINE=99120557; PubMed=9923682;	SEQUENCE FROM N.A. STRAIN=J99:	[2]		"The complete genome sequence of the qastric pathogen Helicobacter	Venter J.C.;	Haves W.S. Borodovsky M. Karon D. Gmith D. Watthey L., Wallin E.,	Berg D.E., Gocayne J.D., Utterback T.R., Peterson J.D., Kelley J.M.	McKenney K., Fitzgerald L.M., Lee N., Adams M.D., Hickey E.K.	Loftus B., Richardson D., Dodson R., Khalak H. G. Glodek A	Nelson K., Quackenbush J., Zhou L., Kirkhess F F Deterson S.	H = B Gill & Boughorter		MEDITUE=00393 / AICC /00392;	SECULENCE FROM N.A.		NCBI_TaxID=210, 85963;	Helicobacter, Grotton Substitution, Helicobacter gloup;	2	Helicobacter bylori (Gambyrobacter bylori 100)	Helicobacter bylor: (Campylobactor prior)	HP0560 OR THP0507	(TrEMBLrel. 14, Last	(TrEMBLrel, 05,	(Tremblrel.	025285; PRELIMINARY; PRT; 26 AA.	860	

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TIGR;
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Gene 245:203-211(2000).
EMBL; AF193251
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
POLYKETIDE SYNTHASE MODULE 6 (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=ATCC15068;
MEDLINE=20179700; PubMed=10713461;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Firmicutes; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Streptomyces venezuelae.
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
POLYKETIDE SYNTHASE MODULE 6 (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=54571;
"Genetic architecture of the polyketide pikromycin series macrolides."; Gene 245:203-211(2000).
EMBL; AF193252; AAF61862.1; ...
NON_TER 30 30
                                                          STRAIN=ATCC19790; MEDLINE=20179700; PubMed=10713461; Xue Y., Wilson D., Sherman D.H.;
                                                                                                                                             Streptomyces narbonensis.
Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
                                                                                                                                                                                                                              6M8760
                                                                                              SEQUENCE FROM N.A.
                                                                                                                     NCBI_TaxID=67333;
                                                                                                                               Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces
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AE001484; AAD06083.1;
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Pred. No. 4e+02;
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Pred. No. 6.3e+02;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mRNA and protein products.";
Gene 38:153-164(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Nucleotide sequence of the immunity region of Bacillus subtilis bacteriophage phi 105: identification of the repressor gene and its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteriophage phi-105.
Viruses; dsDNA viruses, no RNA stage; Tailed phages; Siphoviridae;
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                                                                                                                              Hussey D.J., Nicola M., Moore S., Dobrovic A.;
"The (4:11)(q21:p15) translocation fuses the NUP98 and RAP1GDS1
and is recurrent in T cell acute lymphocytic leukemia.";
submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF133333; AAD54077.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
NUP98-RAP1GDS1 FUSION PROTEIN TYPE 2 (FRAGMENT).
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                                                                 SEQUENCE
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                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                       Eukaryota;
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58.3%;
                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Euteleostomi; primates; Catarrhini; Hominidae; Homo.
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Pred. No. 6.3e+02;
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Pred. No.
                                                                    A68E5D45DDCB205F CRC64;
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O9PS63;
O1-MAY-2000 (TrEMBLrel. 13, Cr.
O1-MAY-2000 (TrEMBLrel. 13, La.
O1-MAY-2000 (TrEMBLrel. 13, La.
O1-MAY-2000 (TREMBLREL. 13, La.
O1-MAY-2000 (TREMBLREL. 13, La.
COLLEGE SA KDA POLYPEPTIDE.
Gallus gallus (Chicken).
Eukaryota: Metazoa, Chordata:
                                                                                                                                                       Bassuk J.A., Berg R.A.;
"A novel 53 kDa polypeptide from chicken embryo.";
J. Biol. Chem. 266:23732-23738(1991).
SEQUENCE 20 AA; 2334 MW; 79B558F295471484 CRC6
                                                                                                                                                                                                           MEDLINE=92084663; PubMed=1660884;
                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                   NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                        Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
GAP JUNCTION SUBUNIT PROTEIN CONNEXIN37 (FRAGMENT).
Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Mouse connexin37: gene structure and promoter analysis."; Biophim. Biophys. Acta 1492:499-504(2000).
EMBL; AF216831; AAF91222.1; -.
NON_TER 23 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=20461860; PubMed=11004519;
Seul K.H., Beyer E.C.;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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nes 6; Conser
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                                                                                                     27.6%;
50.0%;
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                                                                                                   Score 29; DB 13;
Pred. No. 1.9e+03;
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Pred. No. 1.2e+03;
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SEQUENCE.
SEQUENCE-96007483; PubMed=7559503;
MEDLINE=96007483; PubMed=7559503;
Vinay D.S., Raje M., Verma R.K., Mishra G.C.;
Vinaracterization of novel costimulatory molecules. A protein of 38-42
"Characterization of novel costimulatory molecules. A protein of 38-42
"Characterization of novel costimulatory molecules. A protein of 38-42
differentiation.";
J. Biol. Chem. 270:23429-23436(1995).
SEQUENCE 15 AA; 1758 MW; F10C664C976A5D19 CRC64;
                                                                                                                                                                                                                                                                                                                      Q9QUZ5;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
38-42 KDA COSTIMULATORY FACTOR (FRAGMENT).
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                                                                                                                                                                                                                                                                       NCBI_TaxID=10095;
                                                                                                                                                                                                                                                                                 Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eur. J. Bicchem. 267:894-900(2000).
-i- FUNCTION: ANTIBACTERIAL ACTIVITY AGAINST GRAM-POSITIVE BACTERIUM
S.AURBUS. AND GRAM-NEGATIVE BACTERIUM E.COLI. ACTIVE AGAINST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Goraya J., Wang Y., Li Z., O'Flaherty M., Knoop F.C., Platz J.E., Conlon J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolated from the skins of the North American frogs Rana luteiventris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Peptides with antimicrobial activity from four different families
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=20117700; PubMed=10651828;
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Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rana pipiens (Northern leopard frog)
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                              3 LELFENL 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FAMILY.
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                                                            6,
                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27 AA; 2946 MW;
                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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                                                                      26.7%;
85.7%;
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35.0%;
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                                                       Pred. No. 2e+0
0; Mismatches
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Pred. No. 2.
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1FB44EA20187DDA0 CRC64;
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                                                                    2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                            15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27 AA
                                                                                    DB 11;
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                                                                                    Length 15;
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                                                   Gaps
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RESULT
036585
ID 03
AC 03
DT 01
DT 01
DT 01
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OR
OR
OR
OC VI
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sus scrofa (Pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequ
01-MAR-2010 (TrEMBLrel. 16, Last ann
TROPOMYOSIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q9TRA2;
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                                                                                                                                                                                                                                                                                                                                                                                01-JAN-1998 (TrEMBLrel. 05, 01-JAN-1998 (TrEMBLrel. 05, 01-JUN-2000 (TrEMBLrel. 14,
                                                                                                                                                                                                                                                                                                                                                                                                                  036582;
                                01-JAN-1998 (TrEMBLrel.
01-JAN-1998 (TrEMBLrel.
01-JUN-2000 (TrEMBLrel.
ORF-1 (FRAGMENT)
                                                                                                                                                                                                                                                                                  Chatterjee R., Tsarev S., Pilot J., Coursaget P., Purcell R.;
                                                                                                                                                                                                                                                                                                                                                 Hepatitis E virus.
Viruses; ssRNA positive-strand viruses,
                                                                                                                                                                                                                                                                                                                                                                     ORF-1 (FRAGMENT).
Hepatitis E virus.
Viruses; ssRNA positive-strand viruses,
NCBI_TaxID=12461;
                                                                            036585;
                                                                                      036585
                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                        STRAIN-KIRGIZIA (OSH);
                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=12461;
                                                                                                                                                                                Local Similarity 46.2%; tes 6; Conservation
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                                                                                                                                                                 1 PVLELFENLLERL 13
                                                                                                                                           PVLDLTNSILCRV 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                          25 AA; 2772 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                       PRELIMINARY;
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                                             05,
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                       Created)
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                                                                                                                                                                                                                                          5B7D6EC036BBF038 CRC64;
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Pred. No.
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                                                                                                                                                                                                3.2e+03;
                                                                                                                                                                                                                                                                                                                                                    no DNA stage
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               no
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               DNA stage
                                             update)
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RESULT 14
036591
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036588
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Best Local S
Matches 6
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Best Local Similarity 46.2%;
Matches 6; Conservative
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01-JAN-1998 (TrEMBLrel. C
01-JAN-1998 (TrEMBLrel. C
01-JUN-2000 (TrEMBLrel. ORF-1 (FRAGMENT).
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SEQUENCE
                                                                                                                                                                                                                                                                                                           EMBL; AF
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J. Med. Virol. 0:0-0(1997).
EMBL; AF010418; AAB66532.1;
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                                                                                                                         O36591;
O1-JAN-1998 (TrEMBLrel. 05,
O1-JUN-2000 (TrEMBLrel. 14,
                                                                                                                                                                                                                                                                                                                                                Chatterjee R., Tsarev S., Pilot J.,
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. STRAIN=MOROCCO/F13;
                                                                                                                                                                                                                                                                                                                                                                                    Viruses; ssRNA positive-strand viruses, NCBI_TaxID=12461;
                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis E virus.
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                                                                                                      ORF-1 (FRAGMENT).
Hepatitis E virus
                                                                                                                                                                                                                                                                                                  SEQUENCE
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        J. Med. Virol. 0:0-0(1997).
EMBL; AF010420; AAB66538.1;
NON_TER 1 1
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 SEQUENCE
                                    Chatterjee R., Tsarev Purcell R.;
                                                         STRAIN=MOROCCO/F23;
                                                                  SEQUENCE FROM N.A.
                                                                                   Viruses; ssRNA positive-strand viruses, NCBI_TaxID=12461;
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ER 1 1
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  25 AA; 2772 MW;
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                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                24
                                                                                                                                                                                                                                                             26.7%;
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05,
14,
                                                 Pilot
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Pred. No. 3.2e
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             Last sequence up
                                                                                                                           Last sequence update)
Last annotation update)
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                                                                                                                                               Created)
                                                                                                                                                                                                                                                            Score 28; DB 14;
Pred. No. 3.2e+03;
   5B7D6EC036BBF038
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                                                 Coursaget
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                                                                                                no DNA stage
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   CRC64;
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RESULT 15
036594
ID 036594
AC 036594
DT 01-JAN
DT 01-JUN
DT 01-JUN
DT 01-JUN
DT 01-JUN
CC VITUSES
OX NCBI_TE
RN [1]
RP SEQUEN
RA CHATEL
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Search completed: November 20, 2001, 09:46:20 Job time: 174 sec
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Best Local Similarity 46.2
Matches 6; Conservative
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STRAIN=TUNISIA;

Chatterjee R., Tsarev S., Pilot J., Coursaget P., Emerson S., Purcell R.;

J. Med. Virol. 0:0-0(1997).

EMBL; AF010421; AAB66541.1; -.

NON_TER 1 1

NON_TER 25 AA; 2786 MW; B7AD6EC036BBF020 CRC64;
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O36594;
O1-JAN-1998 (TrEMBLrel. 05, Created)
O1-JAN-1998 (TrEMBLrel. 05, Last sequence update)
O1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
ORF-1 (FRAGMENT).
Hepatitis E virus.
Viruses; ssRNA positive-strand viruses, no DNA stage.
NCBI_TaxID=12461;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                       1 PVLELFENLLERL 13
|||:| ::| |:
|12 PVLDLTNSILCRV 24
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| 12 PVLDLTNSILCRV 24
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46.2%;
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Pred. No. 3.2e+03;
4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 14; Length 25;
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